

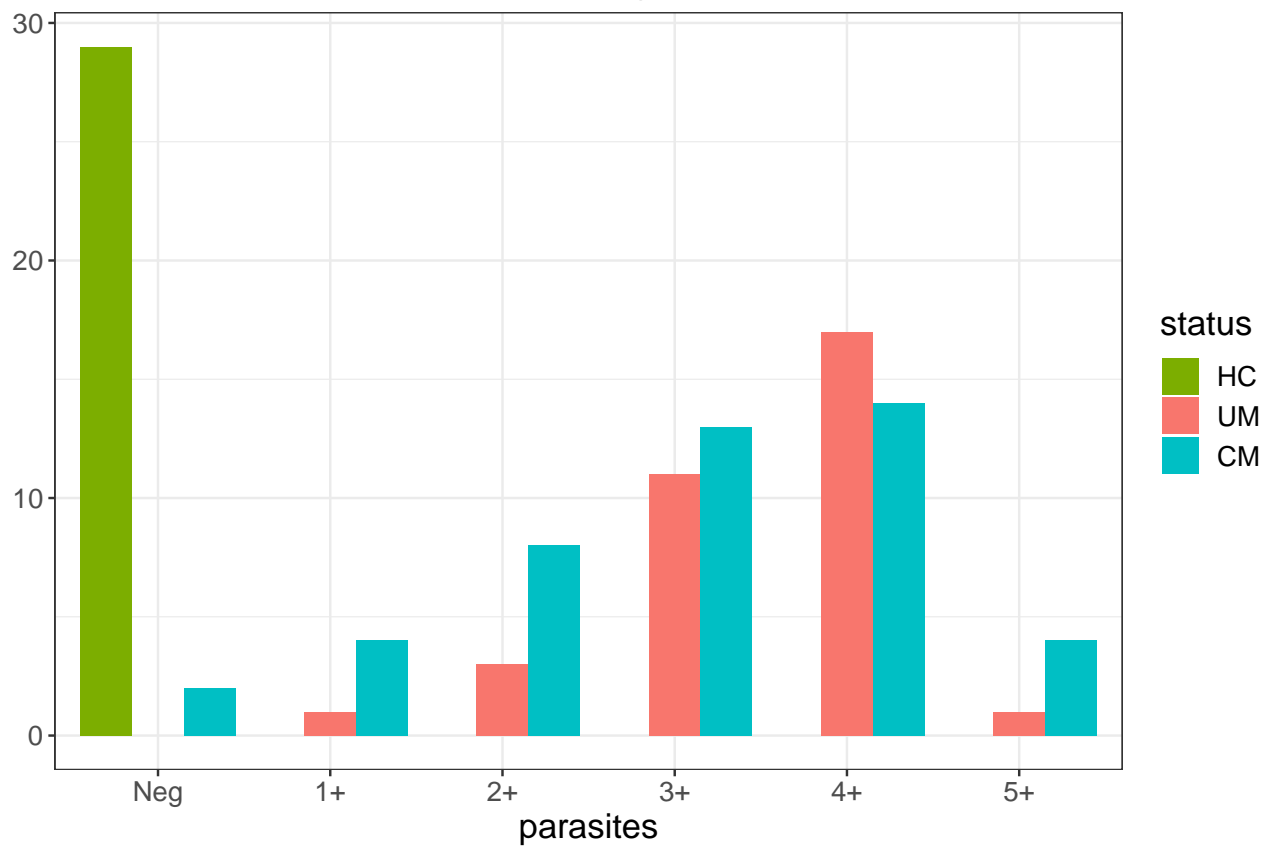
parasitemia

2022-12-18

Parasite counts in each patient group

parasites	HC	UM	CM
Neg	29	0	2
1+	0	1	4
2+	0	3	8
3+	0	11	13
4+	0	17	14
5+	0	1	4

Parasite counts in each patient group



Statistical tests to compare parasites in each patient group

Chi-squared

##

```
## Pearson's Chi-squared test
##
## data: df_parasite_counts %>% dplyr::select(-HC) %>% as.data.frame %>% column_to_rownames("parasite")
## X-squared = 6.6407, df = 5, p-value = 0.2488
```

Ordinal regression (?)

```
## # A tibble: 7 x 6
##   term          coefficient std_error t_stat coef_type      p_val
##   <chr>          <dbl>      <dbl> <dbl> <chr>         <dbl>
## 1 parasites1+    7.45        1.51   4.93 coefficient 0.000000808
## 2 parasites2+    7.05        1.22   5.79 coefficient 0.0000000711
## 3 parasites3+    6.27        1.08   5.81 coefficient 0.0000000621
## 4 parasites4+    5.94        1.05   5.66 coefficient 0.000000148
## 5 parasites5+    7.45        1.51   4.93 coefficient 0.000000808
## 6 HC|UM          2.61        0.733  3.56 scale         0.000377
## 7 UM|CM          6.05        1.02   5.92 scale         0.0000000319
```

Chi-squared (binarize parasites variable)

parasites_bin	UM	CM
low	15	27
high	18	18

```
##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: df_parasite_counts_binned %>% as.data.frame %>% column_to_rownames("parasites_bin")
## X-squared = 1.0883, df = 1, p-value = 0.2968
```

Logistic regression (binarize parasites variable)

```
## # A tibble: 2 x 5
##   term          coefficient std_error t_stat p_value
##   <chr>          <dbl>      <dbl> <dbl> <dbl>
## 1 (Intercept)    0.370      0.241  1.53  0.125
## 2 parasites_binhigh 19.2      1768.  0.0109 0.991
```

HRP & parasitemia in CM patients

hrp2	parasitemia
9531 (5116.5, 10810.5)	29500 (4593, 162200)